

11333US.ST25.txt  
SEQUENCE LISTING

<110> Siegfried Burggraf  
 <120> Method for Detecting Nucleic Acids with internal Control of the Amplification  
 <130> 11333US  
 <150> EP 03015027.0  
 <151> 2003-07-02  
 <150> PCT/EP04/02680  
 <151> 2004-03-15  
 <160> 2  
 <170> PatentIn version 3.1  
 <210> 1  
 <211> 188  
 <212> DNA  
 <213> Hepatitis B virus  
  
 <220>  
 <221> source  
 <222> (1)..(188)  
 <223> GenX  
  
 <400> 1  
 gacgtccttt gtttacgtcc cgtcggcgct gaatcctgcc gacgaccctt ctgggggccg 60  
 cttggggctc tacgccctct ttgccgtctg ccgttccagc caaccacggg gcgcacctct 120  
 ctttacgcgg tctccccgtc tgtgccttct catctgccgg tccgtgtgca cttcgcttca 180  
 cctctgca 188  
  
 <210> 2  
 <211> 120  
 <212> DNA  
 <213> Artificial sequence  
  
 <220>  
 <223> This sequence is an artificial one which only partially coincides with GenX of Hepatitis B (SEQ ID NO:1)  
  
 <220>  
 <221> source  
 <222> (1)..(120)  
 <223> This sequence is an artificial one which only partially coincides with GenX of Hepatitis B (SEQ ID NO:1)  
  
 <400> 2  
 gacgtccttt gtttacgtcc cgtcggcgct gaatcctgcc gacacggggc gcacctctct 60  
 ttacgcggtc taccagtcta tgccttatca tctgcctgtg cacttcgctt cacctctgca 120

## Applicant

-----  
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 PhoneNumber :  
 FaxNumber :  
 EmailAddress :  
 <110> ApplicantName : Siegfried Burggraf

## Application Project

-----  
 <120> Title : Method for Detecting Nucleic Acids with Internal Control of the  
 Amplification  
 <130> AppFileReference : 11333US  
 <140> CurrentAppNumber :  
 <141> CurrentFilingDate : \_\_\_\_-\_\_-\_\_

## Earlier Applications

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 <150> PriorAppNumber : EP 03015027.0  
 <151> PriorFilingDate : 2003-07-03  
 <150> PriorAppNumber : PCT/EP04/02680  
 <151> PriorFilingDate : 2004-03-15

## Sequence

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 <213> OrganismName : Hepatitis B virus  
 <400> PreSequenceString :  
 gacgtccttt gtttacgtcc cgtcggcgct gaatcctgcc gacgaccctt ctcggggccg 60  
 cttggggctc tacgccctct ttgccgtctg ccgttccagc caaccacggg gcgcacctct 120  
 ctttacgcgg tctccccgtc tgtgccttct catctgccgg tccgtgtgca cttcgcttca 180  
 cctctgca 188  
 <212> Type : DNA  
 <211> Length : 188  
 SequenceName : SEQ ID NO:1  
 SequenceDescription :

## Feature

-----  
 Sequence: SEQ ID NO:1:  
 <221> FeatureKey : source  
 <222> LocationFrom : 1  
 <222> LocationTo : 188  
 Other Information : GenX  
 CDSJoin : No

## Sequence

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 <213> OrganismName : Artificial sequence  
 <400> PreSequenceString :  
 gacgtccttt gtttacgtcc cgtcggcgct gaatcctgcc gacacggggc gcacctctct 60  
 ttacgcggtc taccagtcta tgccttatca tctgcctgtg cacttcgctt cacctctgca 120  
 <212> Type : DNA  
 <211> Length : 120  
 SequenceName : SEQ ID NO:2  
 SequenceDescription :

## Feature

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 Sequence: SEQ ID NO:2:  
 <221> FeatureKey : source  
 <222> LocationFrom : 1  
 <222> LocationTo : 120

11333US.workFile.txt

Other Information : This sequence is an artificial one which only partially  
coincides with GenX of Hepatitis B (SEQ ID NO:1)  
CDSJoin : No